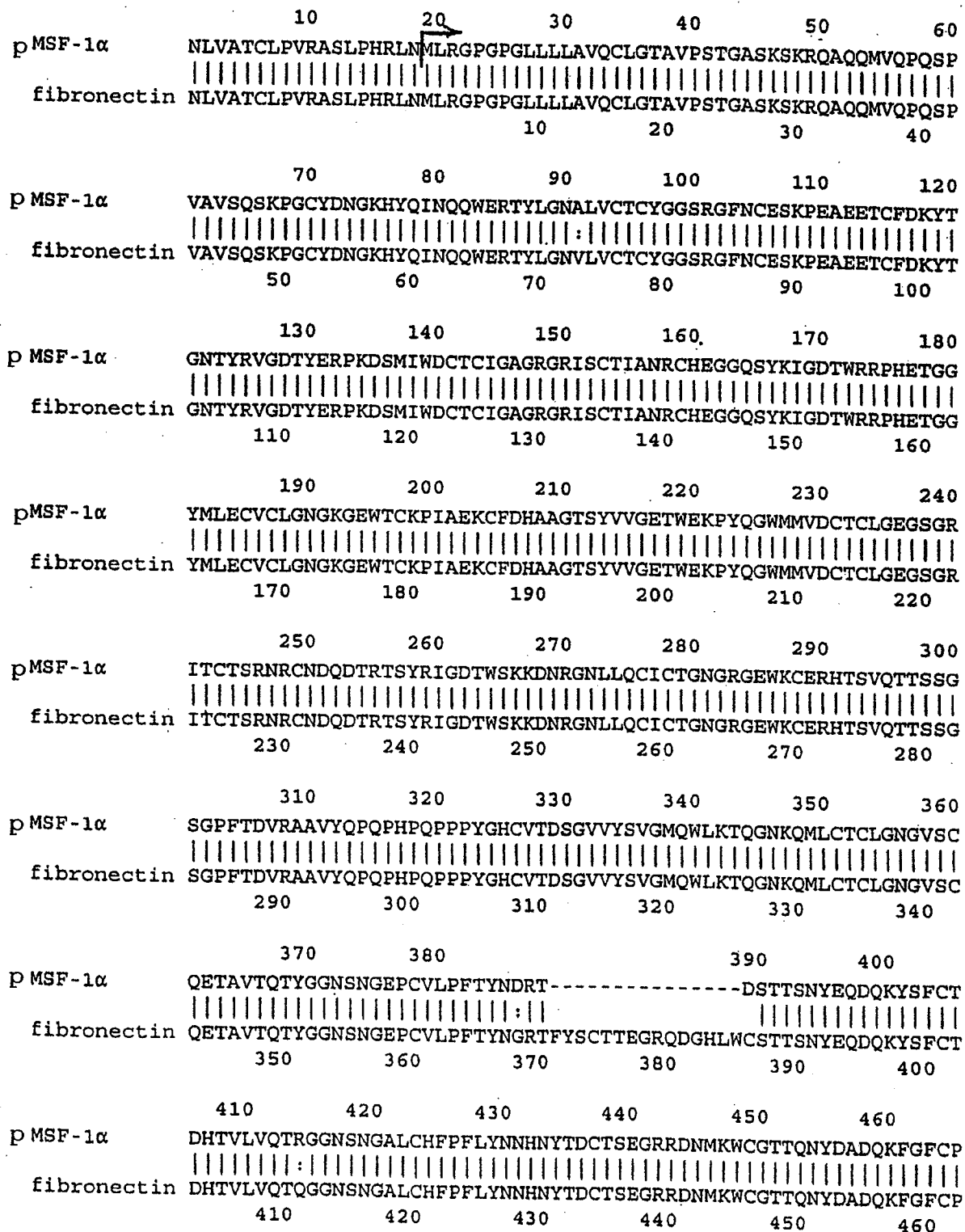


1 CAAACTTGGT GGCAACTTGC CTCCCGGTGC GGGCGTCTCT CCCCCACCGT
51 CTCAA CATGC TTAGGGGTCC GGGGCCCCGGG CTGCTGCTGC TGGCCGTCCA
101 GTGCCTGGGG ACAGCGGTGC CCTCCACGGG AGCCTCGAAG AGCAAGAGGC
151 AGGCTCAGCA AATGGTTCAG CCCCAGTCCC CGGTGGCTGT CAGTCAAAGC
201 AAGCCCCGTT GTTATGACAA TGGAAAACAC TATCAGATAA ATCAACAGTG
251 GGAGCGGACC TACCTAGGCA ATGCGTTGGT TTGTACTTGT TATGGAGGAA
301 GCCGAGGTTT TAACTGCGAG AGTAAACCTG AAGCTGAAGA GACTTGCTTT
351 GACAAGTACA CTGGGAACAC TTACCGAGTG GGTGACACTT ATGAGCGTCC
401 TAAAGACTCC ATGATCTGGG ACTGTACCTG CATCGGGGCT GGGCGAGGGA
451 GAATAAGCTG TACCATCGCA AACCGCTGCC ATGAAGGGGG TCAGTCCTAC
501 AAGATTGGTG ACACCTGGAG GAGACCACAT GAGACTGGTG GTTACATGTT
551 AGAGTGTGTG TGTCTTGGTA ATGGAAAAGG AGAATGGACC TGCAAGCCCA
601 TAGCTGAGAA GTGTTTTGAT CATGCTGCTG GGACTTCCTA TGTGGTCGGA
651 GAAACGTGGG AGAAGCCCTA CCAAGGCTGG ATGATGGTAG ATTGTACTTG
701 CCTGGGAGAA GGCAGCGGAC GCATCACTTG CACTTCTAGA AATAGATGCA
751 ACGATCAGGA CACAAGGACA TCCTATAGAA TTGGAGACAC CTGGAGCAAG
801 AAGGATAATC GAGGAAACCT GCTCCAGTGC ATCTGCACAG GCAACGGCCG
851 AGGAGAGTGG AAGTGTGAGA GGCACACCTC TGTGCAGACC ACATCGAGCG
901 GATCTGGCCC CTTACCGGAT GTTCGTGCAG CTGTTTACCA ACCGCAGCCT
951 CACCCCCAGC CTCCTCCCTA TGGCCACTGT GTCACAGACA GTGGTGTGGT
1001 CTACTCTGTG GGGATGCAGT GGCTGAAGAC ACAAGGAAAT AAGCAAATGC
1051 TTTGCACGTG CCTGGGCAAC GGAGTCAGCT GCCAAGAGAC AGCTGTAACC

Fig. 1 (part 1)

1101 CAGACTTACG GTGGCAACTC AAATGGAGAG CCATGTGTCT TACCATTAC
1151 CTACAACGAC AGGACGGACA GCACAACTTC GAATTATGAG CAGGACCAGA
1201 AATACTCTTT CTGCACAGAC CACACTGTTT TGGTTCAGAC TCGAGGAGGA
1251 AATTCCAATG GTGCCTTGTG CCACTTCCCC TTCCTATACA ACAACCACAA
1301 TTCACTGAT TGCACTTCTG AGGGCAGAAG AGACAACATG AAGTGGTGTG
1351 GGACCACACA GAACTATGAT GCCGACCAGA AGTTTGGGTT CTGCCCCATG
1401 GCTGCCCACG AGGAAATCTG CACAACCAAT GAAGGGGTCA TGTACCGCAT
1451 TGGAGATCAG TGGGATAAGC AGCATGACAT GGGTCACATG ATGAGGTGCA
1501 CGTGTGTTGG GAATGGTCGT GGGGAATGGA CATGCATTGC CTA CTCTCGCAG
1551 CTTCTGAGATC AGTGCATTGT TGATGACATC ACTTACAATG TGAACGACAC
1601 ATTCCACAAG CGTCATGAAG AGGGGCACAT GCTGAACTGT ACATGCTTCG
1651 GTCAGGGTCG GGGCAGGTGG AAGTGTGATC CCGTCGACCA ATGCCAGGAT
1701 TCAGAGACTG GGACGTTTTA TCAAATTGGA GATTCATGGG AGAAGTATGT
1751 GCATGGTGTC AGATACCAGT GCTACTGCTA TGGCCGTGGC ATTGGGGAGT
1801 GGCATTGCCA ACCTTTACAG ACCTATCCAA GCTCAAGTGG TCCTGTGCAA
1851 GTATTTATCA CTGAGACTCC GAGTCAGCCC AACTCCCACC CCATCCAGTG
1901 GAATGCACCA CAGCCATCTC ACATTTCCAA GTACATTCTC AGGTGGAGAC
1951 CTGTGAGTAT CCCACCCAGA AACCTTGGAT ACTGAGTCTC CTAATCTTAT
2001 CAATTCTGAT GGTTTCTTTT TTTCCAGCT TTTGAGCCAA CAACTCTGAT
2051 TAACTATTCC TATAGCATTT ACTATATTG TTTAGTGAAC AAACAATATG
2101 TGGTCAATTA AATTGACTTG TAGACTGAAA AAAAAAAAAA AAAAAA

Fig. 1 (part 2)

*Fig. 2 (part 1)*

	470	480	490	500	510	520
pMSF-1 α	MAAHEEICTTNEGVMYRIGDQWDKQHDMGHMMRCTCVGNRGGEWTCIAYSQLRDQCI VDD					
fibronectin	MAAHEEICTTNEGVMYRIGDQWDKQHDMGHMMRCTCVGNRGGEWTCYAYSQLRDQCI VDD					
	470	480	490	500	510	520
	530	540	550	560	570	580
pMSF-1 α	ITYNVNDTFHKRHEEGHMLNCTCFGQGRGRWKCDPVDQCQDSETGTIFYQIGDSWEKYVHG					
fibronectin	ITYNVNDTFHKRHEEGHMLNCTCFGQGRGRWKCDPVDQCQDSETGTIFYQIGDSWEKYVHG					
	530	540	550	560	570	580
	590	600	610	620	630	640
pMSF-1 α	VRYQCYCYGRGIGEWHCQPLQTYPSSSGPVEVFITETPSQPNSHPIQWNAPQPSHISKYI					
fibronectin	VRYQCYCYGRGIGEWHCQPLQTYPSSSGPVEVFITETPSQPNSHPIQWNAPQPSHISKYI					
	590	600	610	620	630	640
	650	660	670	680	690	700
pMSF-1 α	LRWRPVSI PPRNLGYXVSXSQFXWFLFFPAFEPTTLINYSYSIYYICLVNKQYVVNXID					
	: →					
	(SEQ IS NO.: 18)					
fibronectin	LRWRPKNSVGRWKEATIPGHLNSYTIKGLKPGVVYEGQLISIQYGHQEVTRFDFTTTST					
	650	660	670	680	690	700
	(SEQ IS NO.: 17)					

Fig. 2 (part 2)

SEQ ID NO:	Sequence Type	Binding Site
1	[SEQ ID NO: 13]	5' untranslated region
1	[SEQ ID NO: 14]	Signal
32	[SEQ ID NO: 15]	NH2-terminal segment
52	[SEQ ID NO: 16]	I
97	[SEQ ID NO: 38]	I
141	[SEQ ID NO: 39]	I
186	[SEQ ID NO: 40]	I
232	[SEQ ID NO: 17]	I
273	[SEQ ID NO: 18]	Connecting strand
305	[SEQ ID NO: 19]	I
345	[SEQ ID NO: 20]	II
390	[SEQ ID NO: 28]	II
455	[SEQ ID NO: 21]	I
503	[SEQ ID NO: 22]	I
545	[SEQ ID NO: 23]	I
594	[SEQ ID NO: 24]	III
633	[SEQ ID NO: 25]	Unique Sequence

*VS*SYQF*WLFEPAPFETLLINYSYSIYICLVNKOYVN*IDL*TEKKKKK[SEQ ID NO: 29-33]3' untranslated region

Figure 3

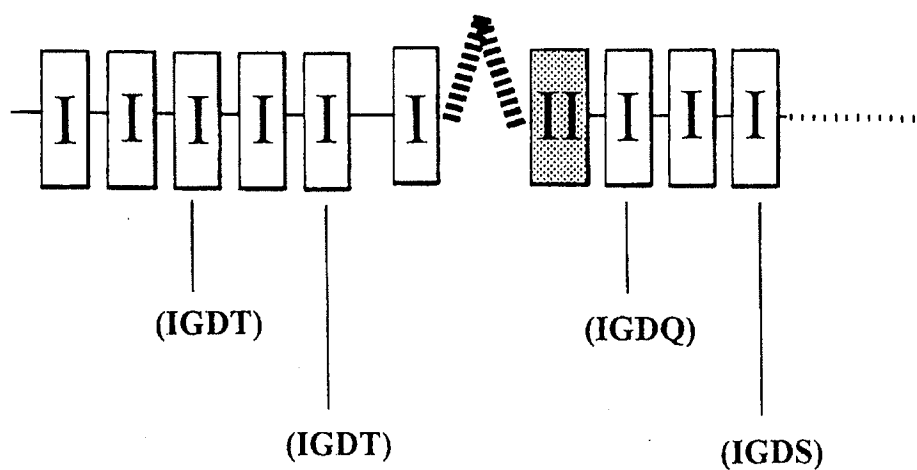


Fig. 4

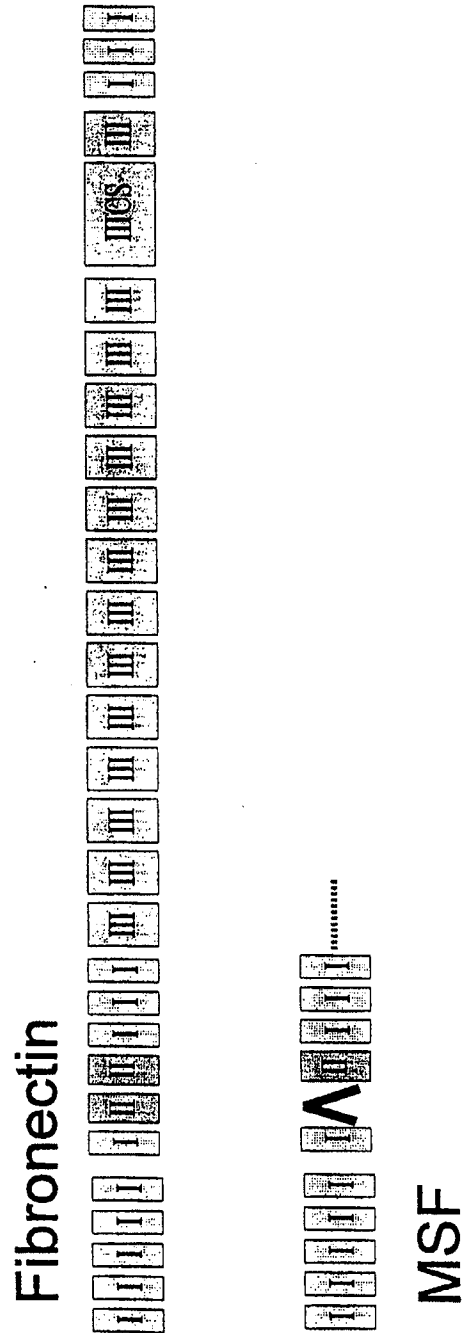


Fig. 5